

AMENDMENTS TO THE SPECIFICATION

Please replace the paragraph beginning at page 5, line 28, with the following amended paragraph:

Figure 1. Amino acid comparison of the intracellular receptor regions (“loops”) of the human Kv1.2 (“hKv1.2”), human Kv1.3 (“hKv1.3”), human Kv1.4 (“hKv1.4”), human Kv1.5 (“hKv1.5”), human Kv1.6 (“hKv1.6”) and human Kv3.4 (“hKv3.4”) are shown in alignment with human Kv1.1 (“hKv1.1”). The black boxes indicate sequence identity; shaded boxes indicate conservative amino acid substitutions. The intracellular receptor regions of hKv1.1, hKv1.2, hKv1.3, hKv1.4, hKv1.5, hKv1.6, and hKv3.4 are designated as hKv1.1LOOP (SEQ ID NO:1), hKv1.2LOOP (SEQ ID NO:1), hKv1.3LOOP (SEQ ID NO:1), hKv1.4LOOP (SEQ ID NO:2), hKv1.5LOOP (SEQ ID NO:24), hKv1.6LOOP (SEQ ID NO:24), and hKv3.4LOOP (SEQ ID NO:25), respectively.

Please replace the paragraph beginning at page 6, line 1, with the following amended paragraph:

Figure 2. Amino acid comparison of the amino-terminal inactivation regions (“N”) of the human Kvβ1b (“hKvβ1b”; also known as “hKvβ1.2”), human Kvβ1c (“hKvβ1c”; also known as “hKvβ1.3”), human Kvβ3 (“hKvβ3”), human Kv1.4 (“hKv1.4”), and human Kv3.4 (“hKv3.4”) are shown in alignment with human Kvβ1 (“hKvβ1”). The black boxes indicate sequence identity; shaded boxes indicate conservative amino acid substitutions. The amino-terminal inactivation regions of hKvβ1, hKvβ1b, hKvβ3, hKv3.4, hKvβ1c, and hKv1.4 are designated as hKvβ1N (SEQ ID NO:5), hKvβ1bN (SEQ ID NO:26), hKvβ3N (SEQ ID NO:26), hKv3.4N (SEQ ID NO:27), hKvβ1cN (SEQ ID NO:28), and hKv1.4N (SEQ ID NO:6), respectively.